

<!--StartFragment-->RESULT 4

AR210003
 LOCUS AR210003 1119 bp DNA linear PAT 20-JUN-2002
 DEFINITION Sequence 1 from patent US 6387637.
 ACCESSION AR210003
 VERSION AR210003.1 GI:21512120
 KEYWORDS .
 SOURCE Unknown.
 ORGANISM Unknown.
 Unclassified.
 REFERENCE 1 (bases 1 to 1119)
 AUTHORS Levin, J.Z., Budziszewski, G.J., Potter, S.L. and Wegrich, L.M.
 TITLE Herbicide target genes and method
 JOURNAL Patent: US 6387637-A 1 14-MAY-2002;
 FEATURES Location/Qualifiers
 source 1..1119
 /organism="unknown"
 /mol_type="unassigned DNA"

ORIGIN

Query Match 89.4%; Score 1100; DB 2; Length 1119;
 Best Local Similarity 99.2%; Pred. No. 0;
 Matches 1119; Conservative 0; Mismatches 0; Indels 9; Gaps 1;

Qy	103	ATGGATGACATGGACACCGTCTACAAGCAATTGGGATTGTTTTCACTAAAGAAGAAGATT	162
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Qy	163	AAAGATGTTGTTCTTAAGGCTGAGATGTTTGACCCGGATGCTCTTGAGCTTGAAGAAGAG	222
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Qy	223	CAGTGGATAAAGCAAGAAGAAACAATGCGTTACTTTGATTATGGGATGATCCCGCTAAA	282
Db	121	CAGTGGATAAAGCAAGAAGAAACAATGCGTTACTTTGATTATGGGATGATCCCGCTAAA	180
Qy	283	TCTGATGAGATTCTTCTCAAATTAGCTGATCGAGCTAAAGCAGTCGATTCCTCAAAGAC	342
Db	181	TCTGATGAGATTCTTCTCAAATTAGCTGATCGAGCTAAAGCAGTCGATTCCTCAAAGAC	240
Qy	343	CTCAAATACAAGGCTGAAGAAGCTAAGCTGATCATACAATTGGGTGAGATGGATGCTATA	402
Db	241	CTCAAATACAAGGCTGAAGAAGCTAAGCTGATCATACAATTGGGTGAGATGGATGCTATA	300
Qy	403	GATTACAGTCTCTTTGAGCAAGCCTATGATTCACTCGATGTAAGTAGATCGTTGCAT	462
Db	301	GATTACAGTCTCTTTGAGCAAGCCTATGATTCACTCGATGTAAGTAGATCGTTGCAT	360
Qy	463	CACTATGAGATGCTAAGCTTCTTAGGGATCAATATGACGCTGAAGCGCTTGTATGATT	522
Db	361	CACTATGAGATGCTAAGCTTCTTAGGGATCAATATGACGCTGAAGCGCTTGTATGATT	420
Qy	523	ATCAAAATCTGGATCTCCAGGCGCAAAATCTCAGGATTTCAGATATGGACAGAGCAAGTT	582
Db	421	ATCAAAATCTGGATCTCCAGGCGCAAAATCT-----CAGATATGGACAGAGCAAGTT	471
Qy	583	GTAAGTATGTATATCAAAATGGGCAGAAAGGCTAGGCCAAAACGCGCGGTGGCTGAGAAA	642
Db	472	GTAAGTATGTATATCAAAATGGGCAGAAAGGCTAGGCCAAAACGCGCGGTGGCTGAGAAA	531
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Db      532  |||||
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Qy      883  ATAGCTGAGAATATGGTTTGTATCCACCATATCCGAGTGGAGTAACACTACAATCTTCA 942
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Db      772  ATAGCTGAGAATATGGTTTGTATCCACCATATCCGAGTGGAGTAACACTACAATCTTCA 831

Qy      943  GGAGAAAGAAACCGGTTTGCAAAACAGGATCAAAGCTCTAAACCGGTTGAAGGCGAAGCTA 1002
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Qy      1003 CTTGTGATAGCAAAAGAGCAAAAGGTTTCGGATGTAAATAAAAATCGACAGCAAGAACATT 1062
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Qy      1063 TTGGAACCGCGGGAAGAAACAGGAGTTATGTCTCTAAGGGTCACAAGATGGTGGTTGAT 1122
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Qy      1123 AGAAAAACCGGTTTGTAGATTCTGGACCTGAAATCGGTCTTGGATGGAACATTGGACCA 1182
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